

Please **replace** Table 3, Page 21 with the following Table 3:

Table 3

Name	Sequence	K _D (μ mol)
ZElan018	K(dns)- STPPSREAYSRPYSVDSDDTNAKHSS HNRRLRTRSRPNG (SEQ ID NO:7)	>50.0
ZElan129	K(dns)-TNAKHSSHNRRLRTR (SEQ ID NO:4)	29.6
ZElan144	K(dns)-rtrlrnhsshkant (SEQ ID NO:1)	28.8
ZElan021	K(dns)- SDHALGTNLRSDNAKEPGDYNCCGNG NSTGRKVFNRRRPSAIP (SEQ ID NO:8)	6.7
ZElan091	K(dns)-PGDYNCCGNGNSTG (SEQ ID NO:6)	0.75
ZElan146	K(dns)-gtsngngccnydgp (SEQ ID NO:3)	21.65

IN THE CLAIMS:

Please **amend** Claim 2 to read as follows:

2. The retro-inverted peptide of claim 1, in which the peptide comprises an amino acid sequence selected from the group consisting of ZElan144 (SEQ ID NO:1), ZElan145 (SEQ ID NO:2) or ZElan146 (SEQ ID NO:3) or a binding "portion thereof."